

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/592,919
Source: IFWP
Date Processed by STIC: 09/25/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>**> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/592,919

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 Invalid Line Length The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.

- 4 Non-ASCII The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for **each** skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

- 8 Skipped Sequences
 (NEW RULES) Sequence(s) 9 missing. If **intentional**, please insert the following lines for **each** skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

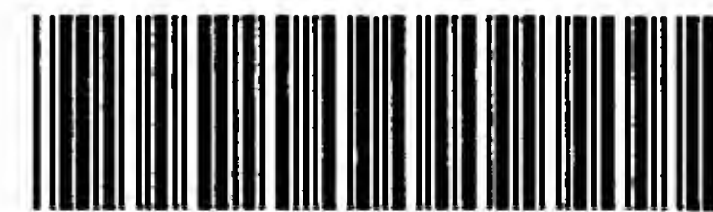
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.

- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown."
 Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 Misuse of n/Xaa "**n**" can **only** represent a single nucleotide; "**Xaa**" can **only** represent a single amino acid



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/592,919

DATE: 09/25/2006

TIME: 10:25:26

Input Set : A:\CORE0037USASEQ.txt

Output Set: N:\CRF4\09252006\J592919.raw

4 <110> APPLICANT: Michael, T. Migawa
 5 Walter F. Lima
 6 Eric E. Swayze
 7 Joshua Nichols
 8 Hongjiang Wu
 9 Thazha P. Prakash
 10 Tadeusz Krzysztof Wyrzykiewicz
 11 Balkrishen Bhat
 12 Stanley T. Crooke
 15 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR OPTIMIZING
 16 CLEAVAGE OF RNA BY RNASE H
 19 <130> FILE REFERENCE: CORE0037USA
 C--> 21 <140> CURRENT APPLICATION NUMBER: US/10/592,919
 C--> 21 <141> CURRENT FILING DATE: 2006-09-15
 21 <150> PRIOR APPLICATION NUMBER: PCT/US2005/008428
 22 <151> PRIOR FILING DATE: 2005-03-15
 24 <150> PRIOR APPLICATION NUMBER: 60/609,516
 25 <151> PRIOR FILING DATE: 2004-09-13
 27 <150> PRIOR APPLICATION NUMBER: 60/567,016
 28 <151> PRIOR FILING DATE: 2004-04-29
 30 <150> PRIOR APPLICATION NUMBER: 60/553,646
 31 <151> PRIOR FILING DATE: 2004-03-15
 33 <160> NUMBER OF SEQ ID NOS: 48
 35 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 37 <210> SEQ ID NO: 1
 38 <211> LENGTH: 20
 39 <212> TYPE: DNA
 40 <213> ORGANISM: Artificial Sequence
 42 <220> FEATURE:
 43 <223> OTHER INFORMATION: Synthetic oligonucleotide
 45 <400> SEQUENCE: 1
 46 ctacgctttc cacgcacagt 20
 48 <210> SEQ ID NO: 2
 49 <211> LENGTH: 20
 50 <212> TYPE: DNA
 51 <213> ORGANISM: Artificial Sequence
 53 <220> FEATURE:
 54 <223> OTHER INFORMATION: Synthetic oligonucleotide
 56 <400> SEQUENCE: 2
 57 agtttaggtc tccgatcgtc 20
 59 <210> SEQ ID NO: 3
 60 <211> LENGTH: 20
 61 <212> TYPE: DNA

Cp8-6)
 Does Not Comply
 Corrected Diskette Needed
 Cp8-3)

RAW SEQUENCE LISTING

DATE: 09/25/2006

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TIME: 10:25:26

Input Set : A:\CORE0037USASEQ.txt

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62 <213> ORGANISM: Artificial Sequence

64 <220> FEATURE:

65 <223> OTHER INFORMATION: Synthetic oligonucleotide

67 <400> SEQUENCE: 3

68 ctgctagcct ctggatttga

20

70 <210> SEQ ID NO: 4

71 <211> LENGTH: 2160

72 <212> TYPE: DNA

73 <213> ORGANISM: Mus musculus

75 <400> SEQUENCE: 4

```

76 ggcgccttgc tctcccgccg gggcgccgga gggggcgggc tggccggcgc acggtgatgt 60
77 ggcgggactc tttgtgcact gcggcaggat acgcgcttgg gcgtcgggac gcggctgcgc 120
78 tcagctctct cctctcgga gctgcagcca tgatggaagt ttgagagttg agccgctgtg 180
79 aggccaggcc cggcgccaggc gagggagatg agagacggcg gcggccacgg cccagagccc 240
80 ctctcagcgc ctgtgagcag ccgcgggggc agcgccctcg gggagccggc cgggcggcgg 300
81 cggcgccaggc ggcggcgggc ctgcgctcct cgctcgtctgt tctaaccggg cagcttctga 360
82 gcagcttcgg agagagacgg tggagaagc cgtgggctcg agcgggagcc ggcgcaggct 420
83 cggcggtctg acctcccgct cctggagcgg gggggagaag cggcgccggc ggccgcggct 480
84 ccggggaggg ggtcggagtc gcctgtcacc attgccaggg ctgggaacgc cggagagttg 540
85 ctctctcccc ttctcctgcc tccaacacgg cggcgccggc ggcggcacgt ccagggaccc 600
86 gggccggtgt taagcctccc gtccgcggcc gccgcacccc ccctggcccg ggctccggag 660
87 gccgcgggag gaggcagccg ctgagaggat tatccgtctt ctccccattc cgctgcctcg 720
88 gctgccaggc ctctggctgc tgaggagaag caggcccagt ctctgcaacc atccagcagc 780
89 cgccgcagca gccattaccc ggctgcggtc cagggccaaag cggcagcaga gcgaggggca 840
90 tcagcgaccg ccaagtccag agccatttcc atcctgcaga agaagcctcg ccaccagcag 900
91 cttctgccat ctctctcctc ctttttcttc agccacaggc tcccagacat gacagccatc 960
92 atcaaagaga tcgttagcag aaacaaaagg agatatcaag aggatggatt cgacttagac 1020
93 ttgacctata ttatccaaa tattattgct atgggatttc ctgcagaaag acttgaaggt 1080
94 gtatacagga acaatattga tgatgtagta aggttttttg attcaaagca taaaaacccat 1140
95 tacaagatat acaatctatg tgctgagaga cattatgaca ccgccaaatt taactgcaga 1200
96 gttgcacagt atccttttga agaccataac ccaccacagc tagaacttat caaacccctc 1260
97 tgtgaagatc ttgaccaatg gctaagtga gatgacaatc atgttgagc aattcactgt 1320
98 aaagctggaa agggacggac tgggtgtaat atttgtgcat atttattgca tcgggggcaa 1380
99 tttttaaagg cacaagaggc cctagatttt tatggggaag taaggaccag agacaaaaag 1440
100 ggagtcacaa ttcccagtc gaggcgctat gtatattatt atagctacct gctaaaaaat 1500
101 cacctggatt acagacccgt ggcactgctg tttcacaaga tgatgtttga aactattcca 1560
102 atgttcagtg gcggaacttg caatcctcag tttgtggtct gccagctaaa ggtgaagata 1620
103 tattcctcca attcaggacc cacgcggcgg gaggacaagt tcatgtactt tgagttccct 1680
104 cagccattgc ctgtgtgtgg tgatatcaaa gtagagttct tccacaaaca gaacaagatg 1740
105 ctcaaaaagg acaaatgtt tcacttttgg gtaaatagc tcttcatacc aggaccagag 1800
106 gaaacctcag aaaaagtgg aaatggaagt ctttgtgatc aggaaatcga tagcatttgc 1860
107 agtatagagc gtgcagataa tgacaaggag tatcttgtac tcaccctaac aaaaaacgat 1920
108 cttgacaaag caacaaaga caaggccaac cgatacttct ctccaaattt taaggtgaaa 1980
109 ctatacttta caaaaacagt agaggagcca tcaaatccag aggctagcag ttcaacttct 2040
110 gtgactccag atgttagtga caatgaacct gatcattata gatattctga caccactgac 2100
111 tctgatccag agaatgaacc ttttgatgaa gatcagcatt cacaattac aaaagtctga 2160

```

114 <210> SEQ ID NO: 5

115 <211> LENGTH: 24

116 <212> TYPE: DNA

RAW SEQUENCE LISTING

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PATENT APPLICATION: US/10/592,919

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Input Set : A:\CORE0037USASEQ.txt

Output Set: N:\CRF4\09252006\J592919.raw

117 <213> ORGANISM: Artificial Sequence
 119 <220> FEATURE:
 120 <223> OTHER INFORMATION: Synthetic oligonucleotide
 122 <400> SEQUENCE: 5
 123 atgacaatca tgttgacgca attc 24
 125 <210> SEQ ID NO: 6
 126 <211> LENGTH: 25
 127 <212> TYPE: DNA
 128 <213> ORGANISM: Artificial Sequence
 130 <220> FEATURE:
 131 <223> OTHER INFORMATION: Synthetic oligonucleotide
 133 <400> SEQUENCE: 6
 134 cgatgcaata aatatgcaca aatca 25
 136 <210> SEQ ID NO: 7
 137 <211> LENGTH: 28
 138 <212> TYPE: DNA
 139 <213> ORGANISM: Artificial Sequence
 141 <220> FEATURE:
 142 <223> OTHER INFORMATION: Synthetic oligonucleotide
 144 <400> SEQUENCE: 7
 145 ctgtaaagct ggaaagggac ggactggt 28
 147 <210> SEQ ID NO: 8
 148 <211> LENGTH: 20
 149 <212> TYPE: DNA
 150 <213> ORGANISM: Artificial Sequence
 152 <220> FEATURE:
 153 <223> OTHER INFORMATION: Synthetic oligonucleotide
 155 <400> SEQUENCE: 8
 156 ccttcctctga aggttcctcc 20
 158 <210> SEQ ID NO: 9
 160 <220> FEATURE:
 161 <223> OTHER INFORMATION: Synthetic oligonucleotide
 163 <400> SEQUENCE: 9
 W--> 164 000
 166 <210> SEQ ID NO: 10
 167 <211> LENGTH: 12
 168 <212> TYPE: RNA
 169 <213> ORGANISM: Artificial Sequence
 171 <220> FEATURE:
 172 <223> OTHER INFORMATION: Synthetic oligonucleotide
 174 <400> SEQUENCE: 10
 175 cgcgaauucg cg 12
 177 <210> SEQ ID NO: 11
 178 <211> LENGTH: 12
 179 <212> TYPE: RNA
 180 <213> ORGANISM: Artificial Sequence
 182 <220> FEATURE:
 183 <223> OTHER INFORMATION: Synthetic oligonucleotide
 185 <400> SEQUENCE: 11

If this is intentionally
 skipped sequence,
 see glen & on error
 summary sheet.

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TIME: 10:25:26

Input Set : A:\CORE0037USASEQ.txt

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```

186 gcgcuaaagc gc 12
188 <210> SEQ ID NO: 12
189 <211> LENGTH: 19
190 <212> TYPE: RNA
191 <213> ORGANISM: Artificial Sequence
193 <220> FEATURE:
194 <223> OTHER INFORMATION: Synthetic oligonucleotide
196 <400> SEQUENCE: 12
197 cgagaggcgg acgggaccg 19
199 <210> SEQ ID NO: 13
200 <211> LENGTH: 21
201 <212> TYPE: DNA
202 <213> ORGANISM: Artificial Sequence
204 <220> FEATURE:
205 <223> OTHER INFORMATION: Synthetic oligonucleotide
207 <220> FEATURE:
208 <221> NAME/KEY: misc_feature
209 <222> LOCATION: 1-19
210 <223> OTHER INFORMATION: Bases at these positions are RNA
212 <400> SEQUENCE: 13
213 cgagaggcgg acgggaccgt t 21
215 <210> SEQ ID NO: 14
216 <211> LENGTH: 21
217 <212> TYPE: DNA
218 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <223> OTHER INFORMATION: Synthetic oligonucleotide
223 <220> FEATURE:
224 <221> NAME/KEY: misc_feature
225 <222> LOCATION: 1-19
226 <223> OTHER INFORMATION: Bases at these positions are RNA
228 <400> SEQUENCE: 14
229 cggtcccgtc cgcctctcgt t 21
231 <210> SEQ ID NO: 15
232 <211> LENGTH: 20
233 <212> TYPE: DNA
234 <213> ORGANISM: Artificial Sequence
236 <220> FEATURE:
237 <223> OTHER INFORMATION: Synthetic oligonucleotide
239 <220> FEATURE:
240 <221> NAME/KEY: misc_feature
241 <222> LOCATION: 4
242 <223> OTHER INFORMATION: N = tetrafluoroindole
244 <400> SEQUENCE: 15
W--> 245 ctgntagcct ctggatttga 20
247 <210> SEQ ID NO: 16
248 <211> LENGTH: 20
249 <212> TYPE: DNA
250 <213> ORGANISM: Artificial Sequence

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RAW SEQUENCE LISTING

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TIME: 10:25:26

Input Set : A:\CORE0037USASEQ.txt

Output Set: N:\CRF4\09252006\J592919.raw

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252 <220> FEATURE:
253 <223> OTHER INFORMATION: Synthetic oligonucleotide
255 <220> FEATURE:
256 <221> NAME/KEY: misc_feature
257 <222> LOCATION: 5
258 <223> OTHER INFORMATION: N = tetrafluoroindole
260 <400> SEQUENCE: 16
W--> 261 ctgcnagcct ctggatttga 20
263 <210> SEQ ID NO: 17
264 <211> LENGTH: 20
265 <212> TYPE: DNA
266 <213> ORGANISM: Artificial Sequence
268 <220> FEATURE:
269 <223> OTHER INFORMATION: Synthetic oligonucleotide
271 <220> FEATURE:
272 <221> NAME/KEY: misc_feature
273 <222> LOCATION: 6
274 <223> OTHER INFORMATION: N = tetrafluoroindole
276 <400> SEQUENCE: 17
W--> 277 ctgctngcct ctggatttga 20
279 <210> SEQ ID NO: 18
280 <211> LENGTH: 20
281 <212> TYPE: DNA
282 <213> ORGANISM: Artificial Sequence
284 <220> FEATURE:
285 <223> OTHER INFORMATION: Synthetic oligonucleotide
287 <220> FEATURE:
288 <221> NAME/KEY: misc_feature
289 <222> LOCATION: 7
290 <223> OTHER INFORMATION: N = tetrafluoroindole
292 <400> SEQUENCE: 18
W--> 293 ctgctancct ctggatttga 20
295 <210> SEQ ID NO: 19
296 <211> LENGTH: 20
297 <212> TYPE: DNA
298 <213> ORGANISM: Artificial Sequence
300 <220> FEATURE:
301 <223> OTHER INFORMATION: Synthetic oligonucleotide
303 <220> FEATURE:
304 <221> NAME/KEY: misc_feature
305 <222> LOCATION: 8
306 <223> OTHER INFORMATION: N = tetrafluoroindole
308 <400> SEQUENCE: 19
W--> 309 ctgctagnct ctggatttga 20
311 <210> SEQ ID NO: 20
312 <211> LENGTH: 20
313 <212> TYPE: DNA
314 <213> ORGANISM: Artificial Sequence
316 <220> FEATURE:

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/592,919

DATE: 09/25/2006
TIME: 10:25:27

Input Set : A:\CORE0037USASEQ.txt
Output Set: N:\CRF4\09252006\J592919.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:15; N Pos. 4
Seq#:16; N Pos. 5
Seq#:17; N Pos. 6
Seq#:18; N Pos. 7
Seq#:19; N Pos. 8
Seq#:20; N Pos. 10
Seq#:21; N Pos. 5
Seq#:22; N Pos. 17
Seq#:23; N Pos. 16
Seq#:24; N Pos. 15
Seq#:25; N Pos. 14
Seq#:26; N Pos. 13
Seq#:27; N Pos. 5,15
Seq#:28; N Pos. 16
Seq#:29; N Pos. 7
Seq#:30; N Pos. 8
Seq#:31; N Pos. 9
Seq#:32; N Pos. 10
Seq#:33; N Pos. 11
Seq#:34; N Pos. 12
Seq#:35; N Pos. 13
Seq#:36; N Pos. 14
Seq#:37; N Pos. 15
Seq#:38; N Pos. 4
Seq#:39; N Pos. 5
Seq#:40; N Pos. 6
Seq#:41; N Pos. 7
Seq#:42; N Pos. 8
Seq#:43; N Pos. 13
Seq#:44; N Pos. 14
Seq#:45; N Pos. 15
Seq#:46; N Pos. 16
Seq#:47; N Pos. 17
Seq#:48; N Pos. 6,16

VERIFICATION SUMMARY

DATE: 09/25/2006

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TIME: 10:25:27

Input Set : A:\CORE0037USASEQ.txt

Output Set: N:\CRF4\09252006\J592919.raw

L:21 M:270 C: Current Application Number differs, Replaced Current Application No

L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:164 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (9) SEQUENCE:

L:245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0

L:261 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0

L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0

L:293 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0

L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0

L:325 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0

L:341 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0

L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0

L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0

L:389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0

L:405 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0

L:421 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0

L:437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0

L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0

L:472 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0

L:491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0

L:510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0

L:527 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0

L:543 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0

L:561 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0

L:577 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0

L:594 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0

L:612 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0

L:628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:0

L:645 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:0

L:662 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:0

L:678 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0

L:694 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:0

L:710 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0

L:726 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:0

L:742 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:0

L:758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 after pos.:0

L:774 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:0

L:790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:0